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RAW SEQUENCE LISTING DATE: 10/31/2001 PATENT APPLICATION: US/09/939,119 TIME: 11:55:39

Input Set : N:\Crf3\RULE60\09939119.txt
Output Set: N:\CRF3\10312001\1939119.raw

SEQUENCE LISTING

```
4 (1) GENERAL INFORMATION:
             (i) APPLICANT: Lipshutz, Robert J.
                             Sapolsky, Ronald
                             Ghandour, Ghassan
      8
     10
            (ii) TITLE OF INVENTION: Polymorphism Detection
           (iii) NUMBER OF SEQUENCES: 14
     12
     14
            (iv) CORRESPONDENCE ADDRESS:
     15
                  (A) ADDRESSEE: Ritter, Van Pelt & Yi LLP
     16
                  (B) STREET: 4906 El Camino Real, Suite 205
     17
                  (C) CITY: Los Altos
     18
                  (D) STATE: CA
                                                                ENTERED
     19
                  (E) COUNTRY: USA
     20
                  (F) ZIP: 94022
     22
             (V) COMPUTER READABLE FORM:
                  (A) MEDIUM TYPE: Floppy disk
     23
                  (B) COMPUTER: IBM PC compatible
     24
     25
                  (C) OPERATING SYSTEM: PC-DOS/MS-DOS
     26
                  (D) SOFTWARE: PatentIn Release #1.0, Version #1.30
     28
            (vi) CURRENT APPLICATION DATA:
C--> 29
                  (A) APPLICATION NUMBER: US/09/939,119
C--> 30
                  (B) FILING DATE: 24-Aug-2001
     31
                  (C) CLASSIFICATION:
     33
           (vii) PRIOR APPLICATION DATA:
     34
                  (A) APPLICATION NUMBER: 08/853,370
     35
                  (B) FILING DATE:
     37
                  (A) APPLICATION NUMBER: 08/563,762
     38
                  (B) FILING DATE: November 29, 1995
     40
          (viii) ATTORNEY/AGENT INFORMATION:
     41
                  (A) NAME: Ritter, Michael J.
     42
                  (B) REGISTRATION NUMBER: 36,653
     43
                  (C) REFERENCE/DOCKET NUMBER: AFFYP016
     45
            (ix) TELECOMMUNICATION INFORMATION:
     46
                  (A) TELEPHONE: 650-903-3500
     47
                  (B) TELEFAX: 650-903-3501
        (2) INFORMATION FOR SEQ ID NO: 1:
             (i) SEQUENCE CHARACTERISTICS:
     52
     53
                  (A) LENGTH: 30 base pairs
     54
                  (B) TYPE: nucleic acid
     55
                  (C) STRANDEDNESS: single
     56
                  (D) TOPOLOGY: linear
            (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 1:
     62 TGCTGCCTTG GTTCRAGCCC TCATCTCTTT
                                                                                 30
       (2) INFORMATION FOR SEQ ID NO: 2:
             (i) SEQUENCE CHARACTERISTICS:
     66
                  (A) LENGTH: 20 base pairs
     67
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(B) TYPE: nucleic acid

68

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```
69
             (C) STRANDEDNESS: single
70
             (D) TOPOLOGY: linear
74
       (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2:
76 AACCAANCTC GGGAGTAGAG
                                                                             20
78 (2) INFORMATION FOR SEQ ID NO: 3:
80
        (i) SEQUENCE CHARACTERISTICS:
81
             (A) LENGTH: 20 base pairs
82
            (B) TYPE: nucleic acid
83
             (C) STRANDEDNESS: single
84
             (D) TOPOLOGY: linear
88
       (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 3:
90 CGGAACCAAN CTCGGGAGTA
                                                                             20
92 (2) INFORMATION FOR SEQ ID NO: 4:
        (i) SEQUENCE CHARACTERISTICS:
95
             (A) LENGTH: 20 base pairs
96
             (B) TYPE: nucleic acid
97
             (C) STRANDEDNESS: single
98
             (D) TOPOLOGY: linear
        (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 4:
104 CGACGGAACC AANCTCGGGA
                                                                              20
106 (2) INFORMATION FOR SEQ ID NO: 5:
108
         (i) SEQUENCE CHARACTERISTICS:
109
              (A) LENGTH: 20 base pairs
110
              (B) TYPE: nucleic acid
111
              (C) STRANDEDNESS: single
              (D) TOPOLOGY: linear
112
116
        (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 5:
118 ACGGAACCAN GTTCGGGAGT
                                                                              20
120 (2) INFORMATION FOR SEQ ID NO: 6:
122
         (i) SEQUENCE CHARACTERISTICS:
123
              (A) LENGTH: 20 base pairs
124
              (B) TYPE: nucleic acid
125
              (C) STRANDEDNESS: single
126
              (D) TOPOLOGY: linear
130
        (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 6:
132 ACGGAACCAN GCTCGGGAGT
                                                                              20
134 (2) INFORMATION FOR SEQ ID NO: 7:
136
         (i) SEQUENCE CHARACTERISTICS:
137
              (A) LENGTH: 20 base pairs
138
              (B) TYPE: nucleic acid
139
              (C) STRANDEDNESS: single
140
              (D) TOPOLOGY: linear
        (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 7:
146 CGGAACCAAN TTCGGGAGTA
                                                                              20
148 (2) INFORMATION FOR SEQ ID NO: 8:
         (i) SEQUENCE CHARACTERISTICS:
150
151
              (A) LENGTH: 20 base pairs
152
              (B) TYPE: nucleic acid
153
              (C) STRANDEDNESS: single
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```
154
               (D) TOPOLOGY: linear
        (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 8:
158
160 CGGAACCAAN CTCGGGAGTA
                                                                              20
162 (2) INFORMATION FOR SEQ ID NO: 9:
         (i) SEQUENCE CHARACTERISTICS:
165
              (A) LENGTH: 20 base pairs
166
              (B) TYPE: nucleic acid
167
               (C) STRANDEDNESS: single
168
               (D) TOPOLOGY: linear
172
        (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 9:
174 GGAACCAAGN TCGGGAGTAG
                                                                              20
176 (2) INFORMATION FOR SEQ ID NO: 10:
178
         (i) SEQUENCE CHARACTERISTICS:
179
              (A) LENGTH: 20 base pairs
180
              (B) TYPE: nucleic acid
181
              (C) STRANDEDNESS: single
182
              (D) TOPOLOGY: linear
186
        (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 10:
188 GAACCAAGTN CGGGAGTAGA
                                                                              20
190 (2) INFORMATION FOR SEQ ID NO: 11:
192
         (i) SEQUENCE CHARACTERISTICS:
193
              (A) LENGTH: 20 base pairs
194
              (B) TYPE: nucleic acid
195
              (C) STRANDEDNESS: single
196
              (D) TOPOLOGY: linear
        (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 11:
202 GAACCAAGCN CGGGAGTAGA
                                                                              20
204 (2) INFORMATION FOR SEQ ID NO: 12:
         (i) SEQUENCE CHARACTERISTICS:
207
              (A) LENGTH: 20 base pairs
208
              (B) TYPE: nucleic acid
209
              (C) STRANDEDNESS: single
210
              (D) TOPOLOGY: linear
214
        (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 12:
216 AACCAAGTTN GGGAGTAGAG
                                                                              20
218 (2) INFORMATION FOR SEQ ID NO: 13:
220
         (i) SEQUENCE CHARACTERISTICS:
              (A) LENGTH: 20 base pairs
221
222
              (B) TYPE: nucleic acid
223
              (C) STRANDEDNESS: single
224
              (D) TOPOLOGY: linear
        (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 13:
228
230 AACCAAGCTN GGGAGTAGAG
                                                                              20
232 (2) INFORMATION FOR SEQ ID NO: 14:
234
         (i) SEQUENCE CHARACTERISTICS:
235
              (A) LENGTH: 30 base pairs
236
              (B) TYPE: nucleic acid
237
              (C) STRANDEDNESS: single
238
              (D) TOPOLOGY: linear
```

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242 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 14:

244 ACCAAAAATC AGTCYGGGTA ACTGAGAGTG

30

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/939,119

DATE: 10/31/2001 TIME: 11:55:40

Input Set : N:\Crf3\RULE60\09939119.txt Output Set: N:\CRF3\10312001\I939119.raw

L:6 M:220 C: Keyword misspelled or invalid format, [(i) APPLICANT:]
L:29 M:220 C: Keyword misspelled or invalid format, [(A) APPLICATION NUMBER:]
L:30 M:220 C: Keyword misspelled or invalid format, [(B) FILING DATE:]